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                 GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on:     September 10, 2004, 20:41:36 ; Search time 94 Seconds
                                          (without alignments)
                                          788.078 Million cell updates/sec

Title:            US-09-919-162F-6
Perfect score:    1244
Sequence:         1 MMPKHCFLGFLISFFLTGVA.........YQPMLDRRSQRSERCVEIP 231

Scoring table:    BLOSUM62
                  Gapop 10.0 , Gapext 0.5

Searched:         1335176 seqs, 320689617 residues

Total number of hits satisfying chosen parameters:      1335176

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :       Published_Applications_AA:*
             1:  /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
             2:  /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
             3:  /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
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             6:  /cgn2_6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pep:*
             7:  /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
             8:  /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
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            11:  /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
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            16:  /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
            17:  /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1244	100.0	231	9	US-09-728-911-2	Sequence 2, Appli
2	1244	100.0	231	9	US-09-949-192-6	Sequence 6, Appli
3	1244	100.0	231	10	US-09-961-404-4	Sequence 4, Appli
4	1244	100.0	231	10	US-09-746-375-33	Sequence 33, Appl
5	1244	100.0	231	12	US-10-245-752-114	Sequence 114, App
6	1244	100.0	231	12	US-10-245-859-114	Sequence 114, App
7	1244	100.0	231	12	US-09-919-162B-6	Sequence 6, Appli
8	1244	100.0	231	14	US-10-245-103-114	Sequence 114, App
9	1244	100.0	231	14	US-10-245-107-114	Sequence 114, App
10	1244	100.0	231	14	US-10-245-143-114	Sequence 114, App
11	1244	100.0	231	14	US-10-245-771-114	Sequence 114, App
12	1244	100.0	231	14	US-10-245-851-114	Sequence 114, App
13	1244	100.0	231	14	US-10-245-883-114	Sequence 114, App
14	1244	100.0	231	14	US-10-237-535-114	Sequence 114, App
15	1244	100.0	231	14	US-10-238-183-114	Sequence 114, App
16	1244	100.0	231	14	US-10-238-283-114	Sequence 114, App
17	1244	100.0	231	14	US-10-238-370-114	Sequence 114, App
18	1244	100.0	231	14	US-10-245-055-114	Sequence 114, App
19	1244	100.0	231	14	US-10-245-147-114	Sequence 114, App
20	1244	100.0	231	14	US-10-245-730-114	Sequence 114, App
21	1244	100.0	231	14	US-10-245-739-114	Sequence 114, App
22	1244	100.0	231	14	US-10-246-210-114	Sequence 114, App
23	1244	100.0	231	14	US-10-239-196-114	Sequence 114, App
24	1244	100.0	231	14	US-10-090-365-2	Sequence 2, Appli
25	1244	100.0	231	14	US-10-243-024-114	Sequence 114, App
26	1244	100.0	231	14	US-10-243-409-114	Sequence 114, App
27	1244	100.0	231	14	US-10-245-621-114	Sequence 114, App
28	1244	100.0	231	14	US-10-245-880-114	Sequence 114, App
29	1244	100.0	231	14	US-10-245-033-114	Sequence 114, App
30	1244	100.0	231	14	US-10-243-095-114	Sequence 114, App
31	1244	100.0	231	14	US-10-245-185-114	Sequence 114, App
32	1244	100.0	231	14	US-10-245-427-114	Sequence 114, App
33	1244	100.0	231	14	US-10-245-473-114	Sequence 114, App
34	1244	100.0	231	14	US-10-245-770-114	Sequence 114, App
35	1244	100.0	231	14	US-10-245-877-114	Sequence 114, App
36	1244	100.0	231	14	US-10-246-976-114	Sequence 114, App
37	1244	100.0	231	14	US-10-243-320-114	Sequence 114, App
38	1244	100.0	231	14	US-10-242-743-114	Sequence 114, App
39	1244	100.0	231	14	US-10-242-845-114	Sequence 114, App
40	1244	100.0	231	14	US-10-104-919-2	Sequence 2, Appli
41	1244	100.0	231	14	US-10-104-919-38	Sequence 38, Appl
42	1244	100.0	231	14	US-10-237-636-114	Sequence 114, App
43	1244	100.0	231	14	US-10-238-325-114	Sequence 114, App
44	1244	100.0	231	14	US-10-238-346-114	Sequence 114, App
45	1244	100.0	231	14	US-10-238-411-114	Sequence 114, App

ALIGNMENTS

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RESULT 1
US-09-728-911-2
;  Sequence 2, Application US/09728911
;  Patent No. US20020012669A1
;  GENERAL INFORMATION:
;  APPLICANT:  Presnell, Scott R.
;  APPLICANT:  Xu, Wenfeng
;  APPLICANT:  Kindsvogel, Wayne
;  APPLICANT:  Chen, Zhi
;  TITLE OF INVENTION:  Human Cytokine Receptor
;  FILE REFERENCE:  99-93
;  CURRENT APPLICATION NUMBER: US/09/728,911
;  CURRENT FILING DATE:  2000-12-01
;  PRIOR APPLICATION NUMBER:  US 60/169,049
;  PRIOR FILING DATE:  1999-12-03
;  PRIOR APPLICATION NUMBER:  US 60/232,219
;  PRIOR FILING DATE:  2000-09-13
;  PRIOR APPLICATION NUMBER:  US 60/244,610
;  PRIOR FILING DATE:  2000-10-31
;  NUMBER OF SEQ ID NOS: 36
;  SOFTWARE:  FastSEQ for Windows Version 3.0
;  SEQ ID NO 2
;    LENGTH: 231
;    TYPE: PRT
;    ORGANISM: Homo sapiens
US-09-728-911-2

  Query Match        100.0%;  Score 1244; DB 9; Length 231;
  Best Local Similarity  100.0%;  Pred. No.2.7e-120;
  Matches 231; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy    1 MMPKHCFLGFLISFFLTGVAGTQSTHESLKPQRVQFQSRNFHNILQWPGRALTGNSSVY 60
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Qy   61 FVQYKIYGQRQWKNKEDCWGTQELSCDLTSETSDIQEPYYGVRAASAGSYSEWSMTPRF 120
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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; CURRENT APPLICATION NUMBER: US/09/961,404
; CURRENT FILING DATE: 2001-09-25
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: PatentIn Ver. 2.1
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; SEQ ID NO 4
; LENGTH: 231
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-961-404-4

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Best Local Similarity   100.0%;  Pred. No. 2.7e-120;
Matches 231; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy    1 MMPKHCFLGFLISFFLTGVAGTQSTHESLKPQRVQFQSRNFHNILQWPGRALTGNSSVY 60
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Db    1 MMPKHCFLGFLISFFLTGVAGTQSTHESLKPQRVQFQSRNFHNILQWPGRALTGNSSVY 60

Qy   61 FVQYKIYGQRQWKNKEDCWGTQELSCDLTSETSDIQEPYYGRVRAASAGSYSEWSMTPRF 120
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Qy  121 TPWWETKIDPPVMNITQVNGSLLVILHAPNLPYRYQEKNVSIEDYYELLYRVFIINNSL 180
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Db  121 TPWWETKIDPPVMNITQVNGSLLVILHAPNLPYRYQEKNVSIEDYYELLYRVFIINNSL 180

Qy  181 EKEQKVYEGAHRAVEIEALTPHSSYCVVAEIYQPMLDRRSQRSEERCVEIP 231
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RESULT 4
US-09-746-375-33
; Sequence 33, Application US/09746375
; Publication No. US20030170823A1
; GENERAL INFORMATION:
; APPLICANT: Presnell, Scott R.
; APPLICANT: Kindsvogel, Wayne
; TITLE OF INVENTION: NOVEL CYTOKINE ZCYTO18
; FILE REFERENCE: 99-106
; CURRENT APPLICATION NUMBER: US/09/746,375
; CURRENT FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 60/172,105
; PRIOR FILING DATE: 1999-12-23
; PRIOR APPLICATION NUMBER: US 60/***,***
; PRIOR FILING DATE: 2000-12-01
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 33
; LENGTH: 231
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-746-375-33

Query Match             100.0%;  Score 1244; DB 10; Length 231;
Best Local Similarity   100.0%;  Pred. No. 2.7e-120;
Matches 231; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy    1 MMPKHCFLGFLISFFLTGVAGTQSTHESLKPQRVQFQSRNFHNILQWPGRALTGNSSVY 60
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Qy   61 FVQYKIYGQRQWKNKEDCWGTQELSCDLTSETSDIQEPYYGRVRAASAGSYSEWSMTPRF 120
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Qy  121 TPWWETKIDPPVMNITQVNGSLLVILHAPNLPYRYQEKNVSIEDYYELLYRVFIINNSL 180
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Qy  181 EKEQKVYEGAHRAVEIEALTPHSSYCVVAEIYQPMLDRRSQRSEERCVEIP 231
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Qy  121 TPWWETKIDPPVMNITQVNGSLLVILHAPNLPYRYQEKNVSIEDYYELLYRVFIINNSL 180
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Qy  181 EKEQKVYEGAHRAVEIEALTPHSSYCVVAEIYQPMLDRRSQRSEERCVEIP 231
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RESULT 2
US-09-949-192-6
; Sequence 6, Application US/09949192
; Patent No. US20020142292A1
; GENERAL INFORMATION:
; APPLICANT: Parhan, Christi L.
; APPLICANT: Gorman, Daniel L.
; APPLICANT: Kurata, Hirokazu
; APPLICANT: Arai, Naoko
; APPLICANT: Sana, Theodore R.
; APPLICANT: Mattson, Jeanine D.
; APPLICANT: Murphy, Erin E.
; APPLICANT: Savkoor, Chetan
; APPLICANT: Grein, Jeffery
; APPLICANT: Smith, Kathleen M.
; APPLICANT: McClanahan, Terrill K.
; TITLE OF INVENTION: MAMMALIAN GENES; RELATED REAGENTS AND METHODS
; FILE REFERENCE: DX011169K
; CURRENT APPLICATION NUMBER: US/09/949,192
; CURRENT FILING DATE: 2001-09-07
; PRIOR APPLICATION NUMBER: 60/231,267
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 53
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 6
; LENGTH: 231
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-949-192-6

Query Match             100.0%;  Score 1244; DB 9; Length 231;
Best Local Similarity   100.0%;  Pred. No. 2.7e-120;
Matches 231; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

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Qy   61 FVQYKIYGQRQWKNKEDCWGTQELSCDLTSETSDIQEPYYGRVRAASAGSYSEWSMTPRF 120
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Qy  121 TPWWETKIDPPVMNITQVNGSLLVILHAPNLPYRYQEKNVSIEDYYELLYRVFIINNSL 180
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Db  181 EKEQKVYEGAHRAVEIEALTPHSSYCVVAEIYQPMLDRRSQRSEERCVEIP 231


RESULT 3
US-09-961-404-4
; Sequence 4, Application US/09961404
; Publication No. US20030022827A1
; GENERAL INFORMATION:
; APPLICANT: WEISS, BERTRAM
; APPLICANT: SABAT, ROBERT
; APPLICANT: ASADULLAH, KHUSRU
; APPLICANT: TOSCHI, LUISELLA
; TITLE OF INVENTION: THREE NEW MEMBERS OF THE CYTOKINE RECEPTOR
; TITLE OF INVENTION: FAMILY CLASS 2
; FILE REFERENCE: SCH-1788
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